Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

- 1. (Currently Amended) Method for the detection in a given DNA sequence of **known and unknown** DNA mutations, single nucleotide polymorphisms, and insertions and deletions comprising the steps of:
- a) producing replicate(s) with an engineered <u>a</u> polymerase of said given DNA sequence with at lest 50% of one of the four natural DNA bases exchanged against a not natural base having at least 50% substitutions in at least one of the four DNA bases;
- b) using said not natural base <u>substitutions</u> to cleave the replicate(s) obtained in step a) and to produce a DNA product presenting sequence-specific fragments;
- c) analyzing said sequence-specific fragments obtained in step b) by mass spectrometry to get sequence-specific fragment patterns; and
- d) using the sequence-specific fragment patterns obtained in step c) to identify sequence changes relative to a reference to said given DNA sequence.
- 2. (Currently Amended) Method according to claim 1 wherein the not natural base substitutions in step a) is are nucleotide equivalents selected from the group consisting of an RNA base (ATP, GTP, CTP, or UTP), a phosphorothioate base, a phosphoroselenoate base, and a photochemically cleavage inducible base.

- 3. (Currently Amended) Method according to claim 1 wherein in the replicate <u>has</u> more than 70% of one of the four natural DNA bases is exchanged against a not natural base substitutions in at least one of the four DNA bases.
- 4. (Currently Amended) Method according to claim 1 wherein in the replicate <u>has</u>
 100% of one of the four natural DNA bases is exchanged against a not natural base
 substitutions in at least one of the four DNA bases.
- 5. (Currently Amended) Method according to claim 2 wherein the RNA base is cleaved in step b) by treatment with alkali and incubation at elevated temperature 55°C.
- 6. (Currently Amended) Method according to claim 2 in which the phosphorothioate or phosphoroselenoate base is cleaved in step b) by condensation of a compound of the nature having the formula OH-(CH2)n-I, where n=2-5, and incubation at elevated temperature 55°C.
- 7. (Original) Method according to claim 2 in which a photochemically cleavage inducible base is cleaved in step b) by exposure to light.
- 8. (Currently Amended) Method according to claim 1 wherein the step a) of producing replicate(s) is carried out with a procedure selected from the group consisting of the polymerase chain reaction OPCR) (PCR) and the linear DNA copying procedure.
- 9. (Original) Method according to claim 8 wherein the linear copying procedure is a rolling circle replication.
- 10. (Previously Presented) Method according to claim 1 comprising further a step a') between step a) and step b), wherein in step a') the replicate(s) is/are purified.

- 11. (Previously Presented) Method according to claim 1 comprising further a step b') between step b) and step c), wherein in step b') the sequence-specific fragments are purified.
- 12. (Currently Amended) Method according to claim 1 wherein the mass spectrometer used for step c) is selected from the group comprising a consisting of MALDI and an ESI mass spectrometers.
- 13. (Currently Amended) Kit for the detection in a given DNA sequence of **known** and unknown DNA mutations, single nucleotide polymorphisms, and insertions and deletions for implementing a method according to claim 1 comprising:

An engineered a DNA polymerase,

A <u>a</u> set of non-natural bases and dNTPs <u>nucleotide equivalents selected from the</u>
group consisting of an RNA base (ATP, GTP, CTP, or UTP), a phosphorothioate base, a

phosphoroselenoate base, and a photochemically cleavage inducible base, and

A a buffer.

- 14. (Previously Presented) Method according to claim 10, wherein the replicate(s) is/are purified on reversed phase material.
- 15. (Currently Amended) Method according to claim 10, wherein the replicate(s) is /are purified with ion exchange resins.
- 16. (Previously Presented) Method according to claim 11, wherein the sequencespecific fragments are purified on reserved phase material.
- 17. (Previously Presented) Method according to claim 11, wherein the sequencespecific fragments are purified with ion exchange resins.